



SEQUENCE LISTING

<110> NAKAMURA, Koji et al.

<120> METHOD OF DETECTING LIVER CANCER, DIAGNOSTIC FOR LIVER
CANCER AND REMEDY FOR CANCER

<130> 0760-0355PUS1

<140> US 10/580,567

<141> 2006-05-26

<150> PCT/JP2004/017499

<151> 2004-11-25

<160> 10

<170> PatentIn version 3.1

<210> 1

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<212> DNA

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<220>

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agccggaccc gcgcccgcac cgctcccggg accgcgaccc cggccgcca gag atg 176
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1acc gcg acc gaa gcc ctc ctg cgc gtc ctc ttg ctc ctg ctg gct ttc 224
Thr Ala Thr Glu Ala Leu Leu Arg Val Leu Leu Leu Leu Leu Ala Phe
5 10 15ggc cac agc acc tat ggg gct gaa tgc ttc ccg gcc tgc aac ccc caa 272
Gly His Ser Thr Tyr Gly Ala Glu Cys Phe Pro Ala Cys Asn Pro Gln
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Asn Gly Phe Cys Glu Asp Asp Asn Val Cys Arg Cys Gln Pro Gly Trp
35 40 45cag ggt ccc ctt tgt gac cag tgc gtg acc tct ccc ggc tgc ctt cac 368
Gln Gly Pro Leu Cys Asp Gln Cys Val Thr Ser Pro Gly Cys Leu His
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gcc aac aac ggg acc tgc gtg agc ctg gac ggt ggc ctc tat gaa tgc	512
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ccc tgt gtg atc aac ggc tcc ccc tgc cag cac gga ggc acc tgc gtg	608
Pro Cys Val Ile Asn Gly Ser Pro Cys Gln His Gly Gly Thr Cys Val	
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Phe Ser Gly Asn Phe Cys Glu Ile Val Ala Asn Ser Cys Thr Pro Asn	
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Pro Cys Glu Asn Asp Gly Val Cys Thr Asp Ile Gly Gly Asp Phe Arg	
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Cys Arg Cys Pro Ala Gly Phe Ile Asp Lys Thr Cys Ser Arg Pro Val	
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Thr Asn Cys Ala Ser Ser Pro Cys Gln Asn Gly Gly Thr Cys Leu Gln	
210 215 220 225	
cac acc cag gtg agc tac gag tgt ctg tgc aag ccc gag ttc aca ggt	896
His Thr Gln Val Ser Tyr Glu Cys Leu Cys Lys Pro Glu Phe Thr Gly	
230 235 240	
ctc acc tgt gtc aag aag cgc gcg ctg agc ccc cag cag gtc acc cgt	944
Leu Thr Cys Val Lys Lys Arg Ala Leu Ser Pro Gln Gln Val Thr Arg	
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ctg ccc agc ggc tat ggg ctg gcc tac cgc ctg acc cct ggg gtg cac	992
Leu Pro Ser Gly Tyr Gly Leu Ala Tyr Arg Leu Thr Pro Gly Val His	
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Glu Leu Pro Val Gln Gln Pro Glu His Arg Ile Leu Lys Val Ser Met	
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Lys Glu Leu Asn Lys Lys Thr Pro Leu Leu Thr Glu Gly Gln Ala Ile	
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Val	Gly	Ile	Val	Phe	Leu	Asn	Lys	Cys	Glu	Thr	Trp	Val	Ser	Asn	Leu	
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cgc	tac	aac	cac	atg	ctg	cgg	aag	aag	aac	ctg	ctg	ctt	cag	tac	aac	1232
Arg	Tyr	Asn	His	Met	Leu	Arg	Lys	Lys	Asn	Leu	Leu	Leu	Gln	Tyr	Asn	
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agc	ggg	gag	gac	ctg	gcc	gtc	aac	atc	atc	ttc	ccc	gag	aag	atc	gac	1280
Ser	Gly	Glu	Asp	Leu	Ala	Val	Asn	Ile	Ile	Phe	Pro	Glu	Lys	Ile	Asp	
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atg	acc	acc	ttc	agc	aag	gag	gcc	ggc	gac	gag	gag	atc	taa			1322
Met	Thr	Thr	Phe	Ser	Lys	Glu	Ala	Gly	Asp	Glu	Glu	Ile				
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ctgtcctaata ctttgtggtg ttcgctatct cttgtgtcaa atctggtgaa cgctacgctt																1442
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His	Gly	Leu	Cys	Gly	Glu	Pro	Gly	Gln	Cys	Ile	Cys	Thr	Asp	Gly	Trp	
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Val Thr Asn Cys Ala Ser Ser Pro Cys Gln Asn Gly Gly Thr Cys Leu
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Arg Leu Pro Ser Gly Tyr Gly Leu Ala Tyr Arg Leu Thr Pro Gly Val
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His Glu Leu Pro Val Gln Gln Pro Glu His Arg Ile Leu Lys Val Ser
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Met Lys Glu Leu Asn Lys Lys Thr Pro Leu Leu Thr Glu Gly Gln Ala
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Thr Val Gly Ile Val Phe Leu Asn Lys Cys Glu Thr Trp Val Ser Asn
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